

GenCore version 4.5  
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OM protein - protein search, using sw model

Run on: August 28, 2002, 17:31:06 ; Search time 75.04 Seconds  
(without alignments)  
312.321 Million cell updates/sec

Title: US-09-502-984B-6

Percent score: 1098  
Sequence: 1 KFEKSKALLAARGPEILCF.....AEPFGGFMSEWSEPSILTR 211

Scoring table:

BLOSUM62  
Gapop 10.0, Gapext 0.5

Searched: 747574 seqs, 111073796 residues

Total number of hits satisfying chosen parameters: 747574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database :

A\_Geneseq\_032802:\*  
1: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1980.DAT:\*  
2: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1981.DAT:\*  
3: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1982.DAT:\*  
4: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1983.DAT:\*  
5: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1984.DAT:\*  
6: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1985.DAT:\*  
7: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1986.DAT:\*  
8: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1987.DAT:\*  
9: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1988.DAT:\*  
10: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1989.DAT:\*  
11: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1990.DAT:\*  
12: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1991.DAT:\*  
13: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1992.DAT:\*  
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15: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1994.DAT:\*  
16: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1995.DAT:\*  
17: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1996.DAT:\*  
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19: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1998.DAT:\*  
20: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA1999.DAT:\*  
21: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA2000.DAT:\*  
22: /SIDSL1/gcgdata/hold-geneseq/geneseqp-emb1/AA2001.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	ID	Description
1	1060	96.5	211	21 AAB21686
2	1060	96.5	225	21 AAB21685
3	1060	96.5	438	21 AA44622
4	1060	96.5	488	18 AAM08349
5	1060	96.5	503	21 AAB13012
6	1060	96.5	508	11 AAR06512
7	1060	96.5	508	16 AAR70032
8	1060	96.5	508	16 AAR69503
9	1053	95.9	508	15 AAR47518
10	1052	95.8	438	21 AA44623
11	869.5	79.2	265	15 AAR50326

12	869.5	79.2	507	11 AAR06511	EPO receptor seque
13	869.5	79.2	507	15 AAR47517	MPL EPO receptor.
14	869.5	79.2	507	16 AAR69502	Mouse erythropoiet
15	862.5	78.6	507	15 AAR50327	Mouse soluble EPO
16	171	15.6	625	22 AAU00377	Mouse thrombopoiet
17	167.5	15.3	117	21 AAY94338	Human cell surface
18	167	15.2	482	16 AAR75941	Soluble murine MPL
19	166	15.1	633	16 AAR79908	Type I MPL recepto
20	166	15.1	633	16 AAR79053	Mouse type I MPL r
21	166	15.1	633	17 AAR98948	Mouse type I MPL r
22	166	15.1	633	17 AAR03513	Mouse type I MPL r
23	166	15.1	633	21 AAY52166	Mouse MPL type I r
24	164	14.9	626	16 AAR75939	Murine myeloprolif
25	160	14.6	30	17 AAR89963	Synthetic human er
26	150	13.7	30	17 AAR89964	Synthetic human er
27	146	13.3	635	13 AAR23970	MPL env protein w
28	146	13.3	635	16 AAR75940	Human myeloprolife
29	146	13.3	635	22 AAU00376	Human thrombopoiet
30	145	13.2	30	17 AAR98936	Synthetic human er
31	143	13.0	30	17 AAR89965	Synthetic human er
32	142	12.9	30	17 AAR98937	Synthetic human er
33	134	12.2	30	17 AAR98938	Synthetic human er
34	127	11.6	32	17 AAR98939	Synthetic human er
35	126	11.5	25	22 AAB48720	Human erythropoiet
36	123.5	11.2	389	20 AAW70846	Human zcytor5 vari
37	122.5	11.2	303	20 AAW70845	Human zcytor5 vari
38	122.5	11.2	350	19 AAW55015	Amino acid sequenc
39	122.5	11.2	350	22 AAE00824	Human NR6 haemopo
40	122.5	11.2	389	20 AAW70848	Human zcytor5 vari
41	122.5	11.2	389	20 AAW70849	Human zcytor5 vari
42	122.5	11.2	389	20 AAW70844	Human zcytor5 vari
43	122.5	11.2	392	20 AAW70840	Human zcytor5 vari
44	122.5	11.2	408	19 AAW59805	Amino acid sequenc
45	122.5	11.2	408	20 AAY26338	Human U4 haematopo

ALIGNMENTS

RESULT 1	
AAB21686	AAB21686 standard; peptide: 211 AA.
ID	
AC	AAB21686;
DT	21-DEC-2000 (first entry)
DE	Human mature erythropoietin receptor EPOR extracellular domain #2.
XX	Ligand; cell surface receptor; erythropoietin; EPOR; human;
KM	protein design automation; PDA.
XX	
OS	Homo sapiens.
XX	
PN	MO200047612-AZ.
PD	17-AUG-2000.
XX	
PF	11-FEB-2000; 2000WO-US03665.
XX	
PR	11-FEB-1999; 99US-0120009.
XX	
PR	29-APR-1999; 99US-0131674.
XX	
PA	(XENC-) XENCOR INC.
XX	
PI	Luo P, Dahiyat B;
XX	
DR	WPI: 2000-549135/50.
XX	
PT	Screening for ligand analogs and agents which modulate ligand-receptor
PT	binding, comprises adding a test ligand to a non-naturally occurring
PT	cell surface receptor analog -
XX	

PS Example 1; Fig 8; 82pp; English.

CC The present invention relates to a method for screening for a ligand  
 CC analog, comprising adding a candidate ligand to a non-naturally occurring  
 CC cell surface receptor analog e.g. erythropoietin receptor (EPOR), and  
 CC determining the binding of the ligand to the analog. The present sequence  
 CC is a mature human erythropoietin receptor (EPOR) extracellular domain.  
 CC Protein Design Automation was carried out on the present sequence, so  
 CC that it may be used in the present invention as a cell surface receptor  
 CC analog.

SO Sequence 211 AA:

Query Match 96.5%; Score 1060; DB 21; Length 211;  
 Best Local Similarity 93.8%; Pred. No. 2e-106;  
 Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFEKSAALLAARGPEELLCTERLEDLVCFEEAASAGVPGNFSPQLEDEPMKLCRL 60  
 DB 1 KFEKSAALLAARGPEELLCTERLEDLVCFEEAASAGVPGNFSPQLEDEPMKLCRL 60  
 QY 61 HOAPTRAGAIRFWCSLPPTADTSSFVPLELRLTAASGAPRHRVTHINEVLLDAPYGLVA 120  
 DB 61 HQAPTRAGAIRFWCSLPPTADTSSFVPLELRLTAASGAPRHRVTHINEVLLDAPYGLVA 120  
 QY 121 RLADSGHVIVRWLPPEPTPMTSHIRFELDISANGAGSVQVRELLEGRTCYLSNLGR 180  
 DB 121 RLADSGHVIVRWLPPEPTPMTSHIRFELDISANGAGSVQVRELLEGRTCYLSNLGR 180  
 QY 181 TRITIAVRARMAEPSPFGFMSAMSEPVSLIT 211  
 DB 181 TRYTFIVRAIRMAEPSPFGFMSAMSEPVSLIT 211

## RESULT 2

AAB21685  
 ID AAB21685 standard; peptide: 225 AA.

AC AAB21685;  
 XX  
 DT 21-DEC-2000 (first entry)

DE Human mature erythropoietin receptor EPOR extracellular domain #1.

XX Ligand; cell surface receptor; erythropoietin; EPOR; human.

OS Homo sapiens.

PN WO200047612-A2.

PD 17-AUG-2000.

PF 11-FEB-2000; 2000WO-US03665.

PR 11-FEB-1999; 99US-0120009.

PR 29-APR-1999; 99US-0131674.

PA (XENC-) XENCOR INC.

PI Luo P, Dahllyat B;

DR WPI: 2000-549135/50.

XX Screening for ligand analogs and agents which modulate ligand-receptor  
 PT binding, comprises adding a test ligand to a non-naturally occurring  
 PT cell surface receptor analog -

PS Example 1; Fig 8; 82pp; English.

CC The present invention relates to a method for screening for a ligand  
 CC analog, comprising adding a candidate ligand to a non-naturally occurring  
 CC cell surface receptor analog e.g. erythropoietin receptor (EPOR), and

CC determining the binding of the ligand to the analog. The present sequence  
 CC is a mature human erythropoietin receptor (EPOR) extracellular domain.  
 CC This sequence may be used in the present invention as a cell surface  
 CC receptor analog.

SO Sequence 225 AA:

Query Match 96.5%; Score 1060; DB 21; Length 225;  
 Best Local Similarity 93.8%; Pred. No. 2.2e-106;  
 Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFEKSAALLAARGPEELLCTERLEDLVCFEEAASAGVPGNFSPQLEDEPMKLCRL 60  
 DB 10 KFEKSAALLAARGPEELLCTERLEDLVCFEEAASAGVPGNFSPQLEDEPMKLCRL 69  
 QY 61 HOAPTRAGAIRFWCSLPPTADTSSFVPLELRLTAASGAPRHRVTHINEVLLDAPYGLVA 120  
 DB 70 HQAPTRAGAIRFWCSLPPTADTSSFVPLELRLTAASGAPRHRVTHINEVLLDAPYGLVA 129  
 QY 121 RLADSGHVIVRWLPPEPTPMTSHIRFELDISANGAGSVQVRELLEGRTCYLSNLGR 180  
 DB 130 RLADSGHVIVRWLPPEPTPMTSHIRFELDISANGAGSVQVRELLEGRTCYLSNLGR 189  
 QY 181 TRITIAVRARMAEPSPFGFMSAMSEPVSLIT 211  
 DB 190 TRYTFIVRAIRMAEPSPFGFMSAMSEPVSLIT 220

## RESULT 3

AA444622  
 ID AA444622 standard; Protein: 438 AA.

AC AA444622;

DT 07-APR-2000 (first entry)

DE Truncated human EPOR(t439).

XX Truncated human EPOR; erythropoietin receptor; hypersensitive EPOR(t439);  
 KW mutant human EPOR; EPOR signalling; cancer; infectious disease; HIV;  
 KW sickle cell anaemia; cytostatic; antitubercular; antiviral;  
 KW immunostimulant; anti-anaemic.

OS Homo sapiens.

PN WO967360-A2.

PD 29-DEC-1999.

PF 25-JUN-1999; 99WO-CA00606.

PR 25-JUN-1998; 98CA-2241576.

PR 25-JAN-1999; 99CA-2260332.

PA (HEMO-) HEMOSOL INC.

PI Bell D, Matthews KE, Mueller SG;

DR WPI: 2000-136979/12.

P-PSDB; AA449634.

XX Serum free defined medium useful for the efficient culture of stem  
 PT cells used for production of hemoglobin -

PS Example 6; Fig 9; 61pp; English.

CC The present sequence is truncated human EPOR (erythropoietin receptor).  
 CC transfection of constitutively active EPOR(t439) by electroporation into  
 CC a cytokine-dependent cell line supports cell population expansion in the  
 CC absence of exogenous cytokines. Mutant human EPOR is used in treatment of  
 CC disorders related to inadequate EPOR signalling. The transfected cells  
 CC may also be used in gene therapy to treat cancer, infectious diseases

CC (e.g. HIV), sickle cell anaemia, and conditions related to abnormal  
 CC expression of erythropoietin.  
 XX  
 SO Sequence 438 AA;

Query Match 96.5%; Score 1060; DB 21; Length 438;  
 Best Local Similarity 93.8%; Pred. No. 5.1e-106;  
 Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFSKALLAARPEELCTERLEDLVCFEEAASAGVPGNFSFQLEDEPMKICRL 60  
 DB 34 kfeskaallaarypeelcterleddlvcfweaasagvpgnysfsyqldepmkicrl 93

QY 61 HOAPTARGAIRFWCSLPTADTSSFVPLELRLTAASGAPRFRHYIHINEVVLDAVGLVA 120  
 DB 94 hgaprtargairfwcslptadtsfvpelrltvaasgaprfrhyihinevvladvglva 153

QY 121 RLADSGHVYIRWLPPPEPTMTSHIRFELDISAGNGAGSVORVELLEGRTCYLSNLGR 180  
 DB 154 rladesghvylrwlppeptmtshirfelveasngagsvqrvellegrtcvlsnlgr 213

QY 181 TRITIAVRARMAEPSPFGFWSANSEVSLIT 211  
 DB 214 trytlavrarmaepsfgfwsawsepsvslit 244

RESULT 4  
 AAM08349  
 ID AAM08349 standard; Protein; 488 AA.  
 AC AAM08349;

XX 14-MAR-1997 (first entry)  
 DE EporFc fusion protein.  
 XX  
 XX Receptor agonist; antibody; erythropoietin receptor; Epor;  
 KM immunogen; antigen; metallochionein; promoter; IgG1; Fc;  
 KW anaemia; therapy.  
 XX  
 OS Chimeric Homo sapiens;  
 OS Chimeric synthetic.  
 XX  
 FH Key Location/Qualifiers  
 FT Domain 1..250  
 FT /label= Epor-ECD  
 FT /note= "erythropoietin receptor extracellular  
 domain"  
 FT Cleavage-site 251..254  
 FT /note= "Factor Xa cleavage site"  
 FT Domain 255..488  
 FT /label= Fc  
 FT /note= "human IgG1 Fc sequence"  
 PN WO9640231-A1.  
 XX  
 PD 19-DEC-1996.  
 XX  
 PF 07-JUN-1996; 96WO-US09613.  
 XX  
 PR 07-JUN-1995; 95US-0474673.  
 XX  
 PA (SMK ) SMITHKLINE BEECHAM CORP.  
 PI Erickson-Miller CL, Young PR;  
 DR WPI; 1997-051900/05.  
 DR N-PSDB; AAT48800.  
 XX  
 PT Recombinant immunogen corresp. to dimeric form of a receptor - used  
 PT for generating antibodies able to act as receptor agonists, esp. of  
 PT erythropoietin receptor for treating anaemia

XX  
 PS Example 1; Page 39-41; 83pp; English.  
 XX  
 CC A fusion protein (AAM08349) encoded by plasmid mta1sEporFc (AAT48800)  
 CC comprises the human erythropoietin receptor (Epor) extracellular  
 CC domain fused (via a Factor Xa cleavage sequence) to the Fc portion  
 CC of human IgG1. It can be expressed e.g. in transfected Drosophila  
 CC S2 cells upon induction with copper sulphate. The cells secrete  
 CC EporFc as a dimeric molecule due to the affinity of the Fc moiety  
 CC for itself. The dimeric receptor can be used as an immunogen to  
 CC generate antibodies (monoclonal, polyclonal, chimeric, humanised)  
 CC able to act as Ecor agonists for use in treatment of anaemia.  
 XX  
 SO Sequence 488 AA;

Query Match 96.5%; Score 1060; DB 18; Length 488;  
 Best Local Similarity 93.8%; Pred. No. 5.9e-106;  
 Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFSKALLAARPEELCTERLEDLVCFEEAASAGVPGNFSFQLEDEPMKICRL 60  
 DB 34 kfeskaallaarypeelcterleddlvcfweaasagvpgnysfsyqldepmkicrl 93

QY 61 HOAPTARGAIRFWCSLPTADTSSFVPLELRLTAASGAPRFRHYIHINEVVLDAVGLVA 120  
 DB 94 hgaprtargairfwcslptadtsfvpelrltvaasgaprfrhyihinevvladvglva 153

QY 121 RLADSGHVYIRWLPPPEPTMTSHIRFELDISAGNGAGSVORVELLEGRTCYLSNLGR 180  
 DB 154 rladesghvylrwlppeptmtshirfelveasngagsvqrvellegrtcvlsnlgr 213

QY 181 TRITIAVRARMAEPSPFGFWSANSEVSLIT 211  
 DB 214 trytlavrarmaepsfgfwsawsepsvslit 244

RESULT 5  
 AAB13012  
 ID AAB13012 standard; Protein; 503 AA.  
 AC AAB13012;

XX 08-DEC-2000 (first entry)  
 DE Q-tagged erythropoietin (Epo) receptor protein.  
 XX  
 XX Site specific label; detection; interaction screening; transglutaminase;  
 KM erythropoietin receptor; Epo.  
 KW  
 XX  
 OS Synthetic.  
 XX  
 PN WO200043492-A2.  
 XX  
 PD 27-JUL-2000.  
 XX  
 PF 20-JAN-2000; 2000WO-US01481.  
 XX  
 PR 22-JAN-1999; 99US-0117327.  
 XX  
 PA (SMK ) SMITHKLINE BEECHAM CORP.  
 PI Tew DG, Powell DJ, Meek TD, Chen W;  
 DR WPI; 2000-499222/44.  
 XX  
 PT Screening for a candidate compound for use in bioassays comprises  
 PT contacting the candidate molecule with a labelled modified protein and  
 PT detecting the label to identify interaction of the two molecules -  
 XX  
 PS Example 4; Page 26; 49pp; English.  
 XX  
 CC This invention relates to methods for the site specific modification of



XX PS Disclosure; Page 27-29; 42pp; English.  
XX XX  
CC The full-length erythropoietin receptor (EPO-R) is given.  
CC Extracellular domains are expressed from vector plasmid pEX-2T as  
CC fusion proteins with glutathione-S-transferase. The domains are  
CC used for investigating the structure of the EPO-R and for  
CC identifying factors involved in regulating differentiation and  
CC proliferation mechanisms in erythroid progenitor cells. They can  
CC also be used for identifying and quantitating EPO and EPO-R as well  
CC as in understanding haematopoietic malignancy and some  
CC cardiovascular system disorders.  
XX CC  
XX CC  
SQ Sequence 508 AA;

Query Match 96.5%; Score 1060; DB 16; Length 508;  
Best Local Similarity 93.8%; Pred. No. 6.2e-106;  
Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFSKAALLAARGPEELLCTERLEDLVCFEEAASAGVGPGRFSPQLEDEPMKLCRL 60  
DB 34 KfskaalllaargpeellcterledlvcfweaasagvpgpnysfsgledepwKLCRL 93  
QY 61 HQAPARGAIRFWCSLPRTADTSSFPVLELRLTAASGAPRFRHYIHINEVLLDAPYGLVA 120  
DB 94 hqapargavrfwcsldpdtssfvpvlelrltvaasgapryhrvihnervlldapvglyva 153  
QY 121 RLADSGHVYIRWLPPEPMTSHIRFELDISAGNGAGSVQRYELLEGRTECVLSNLRGR 180  
DB 154 rladesghvylrwlppecpmtshirfeydvasngagsvqrvellegrtcvlsnlrg 213  
QY 181 TRTIVRARMAEPSPFGFWSAMSEPVSLT 211  
DB 214 trytlavrarmaeppsfgfwswsepsvslt 244

RESULT 8  
AAR69503  
ID AAR69503 standard; Protein; 508 AA.  
XX AAR69503;  
AC AAR69503;  
XX  
XX 11-AUG-1995 (first entry)  
XX DT  
XX DE Human erythropoietin receptor.  
XX KW Erythropoietin receptor; anemia therapy; diagnostic.  
XX KM  
XX OS Homo sapiens.  
XX  
XX Key Location/Qualifiers  
FH 1..24  
FT /note= "signal peptide"  
FT Protein 25..508  
FT /note= "mature protein"  
FT Modified-site 76..79  
FT /note= "N-glycosylation site"  
FT Domain 251..272  
FT /note= "transmembrane region"

US5378808-A.  
XX  
XX 03-JAN-1995.  
XX PD  
XX  
XX 03-FEB-1989; 89US-0306503.  
XX PF  
XX 03-FEB-1989; 89US-0306503.  
XX PR 03-FEB-1989; 89US-0306503.  
XX PR 25-MAR-1991; 91US-0678877.  
XX PR 10-JUN-1993; 93US-0075069.  
XX  
XX (GENV ) GENETICS INST INC.

PI D'andrea A, Jones SS, Wong GC;  
XX  
XX MPI: 1995-051310/07.  
DR N-PSDB; AA081892.  
XX  
XX New recombinant erythropoietin receptor polypeptide(s) - used for  
XX detection, purification, and therapy and for prodn. of antibodies for  
XX detection and therapy  
XX  
XX Claim 2; Fig 9; 24pp; English.  
XX PS  
XX  
XX The sequence is that of a 55-kDa human erythropoietin receptor. The  
XX receptor polypeptide may be used in purification and detection of  
XX erythropoietin, and in production of antibodies for anemia therapy.  
XX The polypeptide may also be used for treating individuals  
XX who are hypersensitive to erythropoietin or who have elevated  
XX erythropoietin levels. They may be used in therapy of e.g. primary  
XX or secondary proliferative polycythemia.  
XX CC  
XX  
SQ Sequence 508 AA;

Query Match 96.5%; Score 1060; DB 16; Length 508;  
Best Local Similarity 93.8%; Pred. No. 6.2e-106;  
Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFSKAALLAARGPEELLCTERLEDLVCFEEAASAGVGPGRFSPQLEDEPMKLCRL 60  
DB 34 KfskaalllaargpeellcterledlvcfweaasagvpgpnysfsgledepwKLCRL 93  
QY 61 HQAPARGAIRFWCSLPRTADTSSFPVLELRLTAASGAPRFRHYIHINEVLLDAPYGLVA 120  
DB 94 hqapargavrfwcsldpdtssfvpvlelrltvaasgapryhrvihnervlldapvglyva 153  
QY 121 RLADSGHVYIRWLPPEPMTSHIRFELDISAGNGAGSVQRYELLEGRTECVLSNLRGR 180  
DB 154 rladesghvylrwlppecpmtshirfeydvasngagsvqrvellegrtcvlsnlrg 213  
QY 181 TRTIVRARMAEPSPFGFWSAMSEPVSLT 211  
DB 214 trytlavrarmaeppsfgfwswsepsvslt 244

RESULT 9  
AAR47518  
ID AAR47518 standard; Protein; 508 AA.  
XX AAR47518;  
AC AAR47518;  
XX  
XX 24-JUN-1994 (first entry)  
XX DT  
XX DE Human EPO receptor.  
XX KW Erythropoietin receptor; recombinant; murine; anaemia.  
XX KM  
XX OS Homo sapiens.  
XX  
XX Key Location/Qualifiers  
FH 1..24  
FT /note= "signal peptide"  
FT Protein 25..508  
FT /note= "mature EPO receptor"  
FT Region 251..272  
FT /note= "putative transmembrane domain"

US5278065-A.  
XX  
XX 11-JAN-1994.  
XX PD  
XX  
XX 03-FEB-1989; 89US-0306503.  
XX PF  
XX 03-FEB-1989; 89US-0306503.  
XX PR 03-FEB-1989; 89US-0306503.  
XX PR 25-MAR-1991; 91US-0678877.  
XX

XX (CHIL-) CHILDRENS MEDICAL CENT.  
 PA (GEM) GENETICS INST INC.  
 PA (WHED) WHITEHEAD INST BIOMEDICAL RES.  
 XX  
 PI D'andrea A, Jones SS, Wong GG;  
 XX  
 DR WPI; 1994-025409/03.  
 DR N-PSDB; AAO53995.  
 XX  
 PT Recombinant DNA encoding erythropoietin receptor - used to  
 PT develop prods. for study, treatment or diagnosis of disorders in  
 PT which receptor is dysfunctional  
 XX  
 PS Disclosure; Fig 9; 24pp; English.  
 XX  
 CC Mouse erythroleukaemia (MEL) cells were used to construct a cDNA  
 CC library. The cDNA was used to transfect COS-1 cells and these were  
 CC screened for radioiodinated erythropoietin (EPO) binding to isolate  
 CC cDNA encoding the EPO receptor. This cDNA was used as a probe to  
 CC screen a human genomic cDNA library to obtain DNA encoding the human  
 CC EPO receptor. The cDNA may be used to study, treat or diagnose  
 CC disorders in which the EPO receptor is dysfunctional. The EPO  
 CC receptor may also be used to raise antibodies or for treating  
 CC hypersensitivity to EPO or who have elevated levels of EPO. The port.  
 CC is pref. used for treating anaemias, primary proliferative polycythemia  
 CC and secondary polycythemia.  
 CC See also AAR47517.  
 CC  
 XX  
 SQ Sequence 508 AA;

Query Match 95.9%; Score 1053; DB 15; Length 508;  
 Best Local Similarity 92.9%; Pred. No. 3.6e-105;

Matches 196; Conservative 12; Mismatches 3; Indels 0; Gaps 0;

OY 1 KFESKAAALNARGPEELCTERLEDVCFEEAASAGVPGNFSFQLEDEPWKLCRL 60  
 |||  
 Db 34 kfeskaallnargpeelcterledvctfweeasagvpgnysfsgyledepwkicrl 93  
 OY 61 HQAPTARGAIRFMCSTPTADTSSFPVLELRITAASGAPRRHRVYHINEVVLDAPVGLVA 120  
 |||  
 Db 94 hqaptargairfmcstptadtsfvpelrtvtaasgaprrhryhinevvlldapvyla 153  
 OY 121 RLADSGHVIRWLPPEPTMTSHIRPELDISAGNGAGSVQRELLGRTCEVLSNLRGR 180  
 |||  
 Db 154 rladesghvirlwlppeptmtshirpeltmshtlyevdvsagngagsvqrveillegrtcevlslnlgr 213  
 OY 181 TRITIAVRARMAEPSPFGFSAMSEPVSLLT 211  
 |||  
 Db 214 trytfavarmaepsfgfwsawsepsyllt 244

# RESULT 10

AAI44623  
 ID AAY44623 standard; Protein; 438 AA.

XX  
 AC AAY44623;

DT 07-APR-2000 (first entry)

DE R154C truncated human EpOR(t439).

XX Truncated human EpOR: erythropoietin receptor; hypersensitive EpOR(t439);

KM mutant human EpOR; EpOR signaling; cancer; infectious disease; HIV;

KW sickle cell anaemia; cytostatic; antimicrobial; antiviral;

XX immunostimulant; anti-anaemic.

XX Homo sapiens.

OS  
 FH Key Location/Qualifiers  
 FT Misc-difference 154 /note- "Wild type Arg substituted by Cys"

XX  
 PN WO967360-A2.  
 XX  
 PD 29-DEC-1999.  
 XX  
 PE 25-JUN-1999; 99MO-CA00606.  
 XX  
 PR 25-JUN-1998; 98CA-2241576.  
 PR 25-JAN-1999; 99CA-2260332.  
 XX  
 PA (HEMO-) HEMOSOL INC.  
 XX  
 PI Bell D, Matthews KE, Mueller SG;  
 XX  
 DR WPI; 2000-136979/12.  
 DR N-PSDB; AA49636.  
 XX  
 PT Serum free defined medium useful for the efficient culture of stem  
 PT cells used for production of hemoglobin -  
 XX  
 PS Example 6; Fig 10; 61pp; English.  
 XX  
 CC The present sequence is R154C truncated human EpOR (erythropoietin  
 CC receptor). Transfection of constitutively active EpOR(t439; R154C) by  
 CC electroporation into a cytokine-dependent cell line supports cell  
 CC population expansion in the absence of exogenous cytokines. Mutant human  
 CC EpOR is used in treatment of disorders related to inadequate EpOR  
 CC signaling. The transfected cells may also used in gene therapy to treat  
 CC cancer, infectious diseases (e.g. HIV), sickle cell anemia, and  
 CC conditions related to abnormal expression of erythropoietin.  
 CC  
 XX  
 SQ Sequence 438 AA;

Query Match 95.8%; Score 1052; DB 21; Length 438;  
 Best Local Similarity 93.4%; Pred. No. 3.8e-105;

Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;

OY 1 KFESKAAALNARGPEELCTERLEDVCFEEAASAGVPGNFSFQLEDEPWKLCRL 60  
 |||  
 Db 34 kfeskaallnargpeelcterledvctfweeasagvpgnysfsgyledepwkicrl 93  
 OY 61 HQAPTARGAIRFMCSTPTADTSSFPVLELRITAASGAPRRHRVYHINEVVLDAPVGLVA 120  
 |||  
 Db 94 hqaptargairfmcstptadtsfvpelrtvtaasgaprrhryhinevvlldapvyla 153  
 OY 121 RLADSGHVIRWLPPEPTMTSHIRPELDISAGNGAGSVQRELLGRTCEVLSNLRGR 180  
 |||  
 Db 154 rladesghvirlwlppeptmtshirpeltmshtlyevdvsagngagsvqrveillegrtcevlslnlgr 213  
 OY 181 TRITIAVRARMAEPSPFGFSAMSEPVSLLT 211  
 |||  
 Db 214 trytfavarmaepsfgfwsawsepsyllt 244

# RESULT 11

AAR50326  
 ID AAR50326 standard; Protein; 265 AA.

XX  
 AC AAR50326;

DT 19-OCT-1994 (first entry)

DE Mouse soluble EPO receptor protein fragment.

XX Murine; soluble; erythropoietin; EPO; receptor protein; sEPO-R; drug;

KW antigen; diagnostic agent; biochemical reagent.

XX Mus musculus.

OS  
 FH Key Location/Qualifiers  
 FT Peptide 1..25 /note- "Signal peptide"

FT Protein 26..265  
 FT /note="Mature Epo-R fragment"  
 XX  
 PN JP06038787-A.  
 XX  
 PD 15-FEB-1994.  
 XX  
 PF 04-MAR-1992; 92JP-0082865.  
 XX  
 PR 04-MAR-1992; . 92JP-0082865.  
 XX  
 PA (SNOW ) SNOW BRAND MILK PROD CO LTD.  
 XX  
 DR WPI; 1994-094847/12.  
 XX  
 DR N-PSDB; AAQ44853.  
 XX  
 PT Soluble erythropoietin receptor protein - and DNA coding for  
 PT SEPO-R, useful as diagnostic reagent  
 XX  
 PS Disclosure; Page 5-6; 9pp; Japanese.  
 XX  
 CC This sequence represents a fragment of the murine soluble erythro-  
 CC poietin (EPO) receptor protein (SEPO-R). This protein is able to  
 CC bind to EPO and has antigenicity as an EPO receptor. The molecular  
 CC weight of the full length protein is pref 33 or 29 kD. The protein  
 CC is useful as a drug, as a diagnostic agent and a biochemical reagent.  
 XX  
 SQ Sequence 265 AA;

Query Match 79.2%; Score 869.5; DB 15; Length 265;  
 Best local similarity 77.7%; Pred. No. 1.1e-85;  
 Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;

QY 1 KFSKAAALLAARPELICTERLEDLVCFEEAASAGVGPNGPNSFOLEDEPMKLCRL 60  
 DB 34 kfskaallasrgseellctqlrledlvctfweaasgm-dfnyfsyqlgesrkscl 92

QY 61 HQAPTARGAIRFWCSLPTADTSSFPLELRFTAASGAPRFRVYIHNEVLLDAPVGLVA 120  
 DB 93 hqaptlrgsvrvcfscslptadtsstfvpelqvtasgspryhrilnhnevlldapaglla 152

QY 121 RLADSGHVIVRMLPPEPTMTSHIRELDISAGNGASVQRYVELLEGRTCYLSNLRGR 180  
 DB 153 rlaeegshvrlwlppegpmtthiryevdvsagngragtgrvlegrtcevlslnrlrg 212

QY 181 TRITIAVRAMAPSPFGGFWSAMSEPSLIT 211  
 DB 213 trytfavarmapsfsgfwswsepsalit 243

RESULT 12  
 ID AAR06511 standard; protein; 507 AA.  
 AC AAR06511;  
 XX  
 DT 04-JUN-1991 (first entry)  
 XX  
 DE EPO receptor sequence deduced from DNA of clone 190.  
 XX  
 KM Erythropoietin; Diamond Blackfan anaemia; polycythemia vera.  
 XX  
 OS Mus musculus.  
 XX  
 FH Key Location/Qualifiers  
 FT Peptide 1..24  
 FT /label=signal peptide  
 FT Domain 25..248  
 FT /label=extracellular domain  
 FT /note=EPO binding region  
 FT 248..271  
 FT /label=transmembrane domain

FT Domain 272..507  
 FT /label=intracellular domain  
 FT Modified-site 75..77  
 FT /label=N-linked\_glycos  
 FT Modified-site 182..184  
 FT /label=N-linked\_glycos  
 PN WO9008822-A.  
 XX  
 PD 09-AUG-1990.  
 XX  
 PF 01-FEB-1990; 90WO-US00635.  
 XX  
 PR 03-FEB-1989; 89US-0306503.  
 XX  
 PA (GENE-) GENETICS INST INC.  
 PA (WHIT-) WHITEHEAD INST.  
 XX  
 PI D'andrea A, Mong G;  
 XX  
 DR WPI; 1990-260931/34.  
 XX  
 DR N-PSDB; AAQ05747.  
 XX  
 PT Erythropoietin receptor and gene - used for developing reagents  
 PT and systems to control and study erythropoiesis.  
 XX  
 PS Disclosure; Fig 1; 53pp; English.  
 XX  
 CC The sequence was deduced from DNA from a clone isolated from a  
 CC CDNA library prep. from uninduced murine erythroleukemia cells.  
 CC It is a type I transmembrane protein with binding affinity for EPO.  
 CC The gene and recombinant EPO receptor produced on expression of  
 CC the DNA are used to develop reagents and systems to control and  
 CC study erythropoiesis. It is believed that the EPO receptor is  
 CC dysfunctional in individuals with Diamond Blackfan anaemia, and  
 CC may be hyperactive in polycythemia vera.  
 CC See also AAR06512 (human EPO receptor).  
 XX  
 SQ Sequence 507 AA;

Query Match 79.2%; Score 869.5; DB 11; Length 507;  
 Best local similarity 77.7%; Pred. No. 2.5e-85;  
 Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;

QY 1 KFSKAAALLAARPELICTERLEDLVCFEEAASAGVGPNGPNSFOLEDEPMKLCRL 60  
 DB 34 kfskaallasrgseellctqlrledlvctfweaasgm-dfnyfsyqlgesrkscl 92

QY 61 HQAPTARGAIRFWCSLPTADTSSFPLELRFTAASGAPRFRVYIHNEVLLDAPVGLVA 120  
 DB 93 hqaptlrgsvrvcfscslptadtsstfvpelqvtasgspryhrilnhnevlldapaglla 152

QY 121 RLADSGHVIVRMLPPEPTMTSHIRELDISAGNGASVQRYVELLEGRTCYLSNLRGR 180  
 DB 153 rlaeegshvrlwlppegpmtthiryevdvsagngragtgrvlegrtcevlslnrlrg 212

QY 181 TRITIAVRAMAPSPFGGFWSAMSEPSLIT 211  
 DB 213 trytfavarmapsfsgfwswsepsalit 243

RESULT 13  
 ID AAR47517 standard; protein; 507 AA.  
 AC AAR47517;  
 XX  
 DT 24-JUN-1994 (first entry)  
 XX  
 DE MEL EPO receptor.  
 XX  
 KM Erythropoietin receptor; recombinant; murine; anaemia.







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